SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Keller, Gordon M. Kennedy, Marion Choi, Kyunghee Firpo, Meri T.
- (ii) TITLE OF INVENTION: NOVEL EMBRYONIC CELL POPULATIONS AND METHODS TO ISOLATE SUCH POPULATIONS
- (iii) NUMBER OF SEQUENCES: 4
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Sheridan Ross & McIntosh
 - (B) STREET: 1700 Lincoln St., Suite 3500
 - (C) CITY: Denver
 - (D) STATE: Colorado
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 80203
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk

 - (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

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- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/343,686
 - (B) FILING DATE: 21-NOV-1994
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Kovarik, Joseph E.
 - (B) REGISTRATION NUMBER: 33,005
 - (C) REFERENCE/DOCKET NUMBER: 2879-26
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 303/863-9700
 - (B) TELEFAX: 303/863-0223





(2) INFORMATION FOR SEQ ID NO:1:
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear
(ii) MOLECULE TYPE: DNA (genomic)
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
TGGTGGAGTC TGGGGGAGGC TTA
(2) INFORMATION FOR SEQ ID NO:2:
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear
(ii) MOLECULE TYPE: DNA (genomic)
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
GGCTCCCTCA GGGACAAATA TCCA
(2) INFORMATION FOR SEQ ID NO:3:
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 32 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: DNA (genomic)
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
CGCGGCCCCA AGCTTGTTAA CATCGATGGA TG
(2) INFORMATION FOR SEQ ID NO:4:
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear
<pre>(ii) MOLECULE TYPE: DNA (genomic)</pre>

(xi) SEQUENCE DESCRIPTION: SEQ ID

GGCGTTACTT AAGCTAGCTT GCCAAAGGTA C

NO:1:			23
NO:2:		.or	24
NO:3:			32
NO:4:			31

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